

FIGURE 1

ACCCACGCGC AGCGGCCGGA GATGCAGCGG GGCGCCGCGC TGTGCCTGCG ACTGTGGCTC TGCCTGGGAC TCCTGGACGG CCTGGTGAGT GACTACTCCA TGACCCCCC GACCTTGAAC ATCACGGAGG AGTCACACGT CATCGACACC GGTGACAGCC TGTCCATCTC CTGCAGGGGA CAGCACCCC TCGAGTGGC TTGGCCAGGA GCTCAGGAGG CGCCAGCCAC CGGAGACAAG GACAGCGAGG ACACGGGGGT GGTGCGAGAC TGCGAGGGCA CAGACGCCAG GCCCTACTGC AAGGTGTTGC TGCTGCACGA GGTACATGCC AACGACACAG GCAGCTACGT CTGCTACTAC AAGTACATCA AGGCACGCAT CGAGGGCACC ACGGCCGCCA GCTCCTACGT GTTCGTGAGA 421 GACTTTGAGC AGCCATTCAT CAACAAGCCT GACACGCTCT TGGTCAACAG GAAGGACGCC ATGTGGGTGC CCTGTCTGGT GTCCATCCCC GGCCTCAATG TCACGCTGCG CTCGCAAAGC TCGGTGCTGT GGCCAGACGG GCAGGAGGTG GTGTGGGATG ACCGGCGGGG CATGCTCGTG TCCACGCCAC TGCTGCACGA TGCCCTGTAC CTGCAGTGCG AGACCACCTG GGGAGACCAG GACTTCCTTT CCAACCCCTT CCTGGTGCAC ATCACAGGCA ACGAGCTCTA TGACATCCAG 661 CTGTTGCCCA GGAGTCGCT GGAGCTGCTG GTAGGGGAGA AGCTGGTCCT CAACTGCACC GTGTGGGCTG AGTTTAACTC AGGTGTCACC TTTGACTGGG ACTACCCAGG GAAGCAGGCA GAGCGGGGTA AGTGGGTGCC CGAGCGACGC TCCCAACAGA CCCACACAGA ACTCTCCAGC ATCCTGACCA TCCACAACGT CAGCCAGCAC GACCTGGGCT CGTATGTGTG CAAGGCCAAC AACGCCATCC AGCGATTTCG GGAGAGCACC GAGGTCATTG TGCATGAAAA TCCCTTCATC 1021 AGCGTCGAGT GGCTCAAAGG ACCCATCCTG GAGGCCACGG CAGGAGACGA GCTGGTGAAG 1081 CTGCCCGTGA AGCTGGCAGC GTACCCCCCG CCCGAGTTCC AGTGGTACAA GGATGGAAAG 1141 GCACTGTCCG GGCGCCACAG TCCACATGCC CTGGTGCTCA AGGAGGTGAC AGAGGCCAGC 1201 ACAGGCACCT ACACCCTCGC CCTGTGGAAC TCCGCTGCTG GCCTGAGGCG CAACATCAGC 1261 CTGGAGCTGG TGGTGAATGT GCCCCCCAG ATACATGAGA AGGAGGCCTC CTCCCCCAGC 1321 ATCTACTCGC GTCACAGCCG CCAGGCCCTC ACCTGCACGG CCTACGGGGT GCCCCTGCCT 1381 CTCAGCATCC AGTGGCACTG GCGGCCCTGG ACACCCTGCA AGATGTTTGC CCAGCGTAGT 1441 CTCCGGCGGC GGCAGCAGCA AGACCTCATG CCACAGTGCC GTGACTGGAG GGCGGTGACC 1501 ACGCAGGATG CCGTGAACCC CATCGAGAGC CTGGACACCT GGACCGAGTT TGTGGAGGGA 1561 AAGAATAAGA CTGTGAGCAA GCTGGTGATC CAGAATGCCA ACGTGTCTGC CATGTACAAG 1621 TGTGTGGTCT CCAACAAGGT GGGCCAGGAT GAGCGGCTCA TCTACTTCTA TGTGACCACC 1681 ATCCCCGACG GCTTCACCAT CGAATCCAAG CCATCCGAGG AGCTACTAGA GGGCCAGCCG 1741 GTGCTCCTGA GCTGCCAAGC CGACAGCTAC AAGTACGAGC ATCTGCGCTG GTACCGCCTC 1801 AACCTGTCCA CGCTGCACGA TGCGCACGGG AACCCGCTTC TGCTCGACTG CAAGAACGTG 1861 CATCTGTTCG CCACCCCTCT GGCCGCCAGC CTGGAGGAGG TGGCACCTGG GGCGCCAC 1921 GCCACGCTCA GCCTGAGTAT CCCCCGCGTC GCGCCCGAGC ACGAGGGCCA CTATGTGTGC 1981 GAAGTGCAAG ACCGGCGCAG CCATGACAAG CACTGCCACA AGAAGTACCT GTCGGTGCAG 2041 GCCCTGGAAG CCCCTCGGCT CACGCAGAAC TTGACCGACC TCCTGGTGAA CGTGAGCGAC 2101 TCGCTGGAGA TGCAGTGCTT GGTGGCCGGA GCGCACGCGC CCAGCATCGT GTGGTACAAA 2161 GACGAGAGGC TGCTGGAGGA AAAGTCTGGA GTCGACTTGG CGGACTCCAA CCAGAAGCTG 2221 AGCATCCAGC GCGTGCGCGA GGAGGATGCG GGACCGTATC TGTGCAGCGT GTGCAGACCC 2281 AAGGGCTGCG TCAACTCCTC CGCCAGCGTG GCCGTGGAAG GCTCCGAGGA TAAGGGCAGC 2341 ATGGAGATCG TGATCCTTGT CGGTACCGGC GTCATCGCTG TCTTCTTCTG GGTCCTCCTC 2401 CTCCTCATCT TCTGTAACAT GAGGAGGCCG GCCCACGCAG ACATCAAGAC GGGCTACCTG 2461 TCCATCATCA TGGACCCCGG GGAGGTGCCT CTGGAGGAGC AATGCGAATA CCTGTCCTAC 2521 GATGCCAGCC AGTGGGAATT CCCCCGAGAG CGGCTGCACC TGGGGAGAGT GCTCGGCTAC 2581 GGCGCCTTCG GGAAGGTGGT GGAAGCCTCC GCTTTCGGCA TCCACAAGGG CAGCAGCTGT 2641 GACACCGTGG CCGTGAAAAT GCTGAAAGAG GGCGCCACGG CCAGCGAGCA GCGCGCGCTG 2701 ATGTCGGAGC TCAAGATCCT CATTCACATC GGCAACCACC TCAACGTGGT CAACCTCCTC 2761 GGGGCGTGCA CCAAGCCGCA GGGCCCCCTC ATGGTGATCG TGGAGTTCTG CAAGTACGGC 2821 AACCTCTCCA ACTTCCTGCG CGCCAAGCGG GACGCCTTCA GCCCCTGCGC GGAGAAGTCT 2881 CCCGAGCAGC GCGGACGCTT CCGCGCCATG GTGGAGCTCG CCAGGCTGGA TCGGAGGCGG 2941 CCGGGGAGCA GCGACAGGGT CCTCTTCGCG CGGTTCTCGA AGACCGAGGG CGGAGCGAGG 3001 CGGGCTTCTC CAGACCAAGA AGCTGAGGAC CTGTGGCTGA GCCCGCTGAC CATGGAAGAT 3061 CTTGTCTGCT ACAGCTTCCA GGTGGCCAGA GGGATGGAGT TCCTGGCTTC CCGAAAGTGC 3121 ATCCACAGAG ACCTGGCTGC TCGGAACATT CTGCTGTCGG AAAGCGACGT GGTGAAGATC 3181 TGTGACTTTG GCCTTGCCCG GGACATCTAC AAAGACCCCG ACTACGTCCG CAAGGGCAGT



| 3241 | GCCCGGCTGC | CCCTGAAGTG | GATGGCCCCT | GAAAGCATCT | TCGACAAGGT | GTACACCACG |
|------|------------|------------|------------|------------|------------|------------|
| 3301 | CAGAGTGACG | TGTGGTCCTT | TGGGGTGCTT | CTCTGGGAGA | TCTTCTCTCT | GGGGGCCTCC |
| 3361 | CCGTACCCTG | GGGTGCAGAT | CAATGAGGAG | TTCTGCCAGC | GCGTGAGAGA | CGGCACAAGG |
| 3421 | ATGAGGGCCC | CGGAGCTGGC | CACTCCCGCC | ATACGCCACA | TCATGCTGAA | CTGCTGGTCC |
| 3481 | GGAGACCCCA | AGGCGAGACC | TGCATTCTCG | GACCTGGTGG | AGATCCTGGG | GGACCTGCTC |
| 3541 | CAGGGCAGGG | GCCTGCAAGA | GGAAGAGGAG | GTCTGCATGG | CCCCGCGCAG | CTCTCAGAGC |
| 3601 | TCAGAAGAGG | GCAGCTTCTC | GCAGGTGTCC | ACCATGGCCC | TACACATCGC | CCAGGCTGAC |
| 3661 | GCTGAGGACA | GCCCGCCAAG | CCTGCAGCGC | CACAGCCTGG | CCGCCAGGTA | TTACAACTGG |
| 3721 | GTGTCCTTTC | CCGGGTGCCT | GGCCAGAGGG | GCTGAGACCC | GTGGTTCCTC | CAGGATGAAG |
| 3781 | ACATTTGAGG | AATTCCCCAT | GACCCCAACG | ACCTACAAAG | GCTCTGTGGA | CAACCAGACA |
| 3841 | GACAGTGGGA | TGGTGCTGGC | CTCGGAGGAG | TTTGAGCAGA | TAGAGAGCAG | GCATAGACAA |
| 3901 | GAAAGCGGCT | TCAGCTGTAA | AGGACCTGGC | CAGAATGTGG | CTGTGACCAG | GGCACACCCT |
| 3961 | GACTCCCAAG | GGAGGCGGCG | GCGGCCTGAG | CGGGGGCCC | GAGGAGGCCA | GGTGTTTTAC |
| 4021 | AACAGCGAGT | ATGGGGAGCT | GTCGGAGCCA | AGCGAGGAGG | ACCACTGCTC | CCCGTCTGCC |
| 4081 | CGCGTGACTT | TCTTCACAGA | CAACAGCTAC | TAA | | |
| | | | | | | |

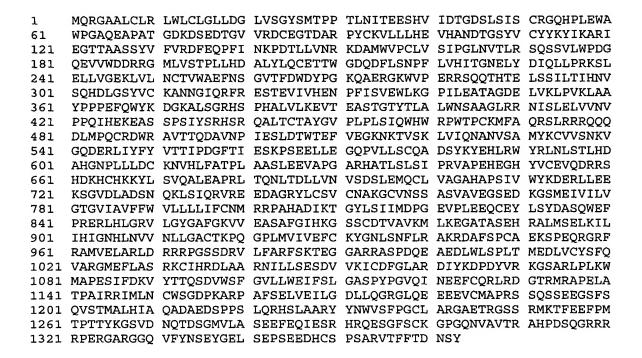


FIGURE 2B

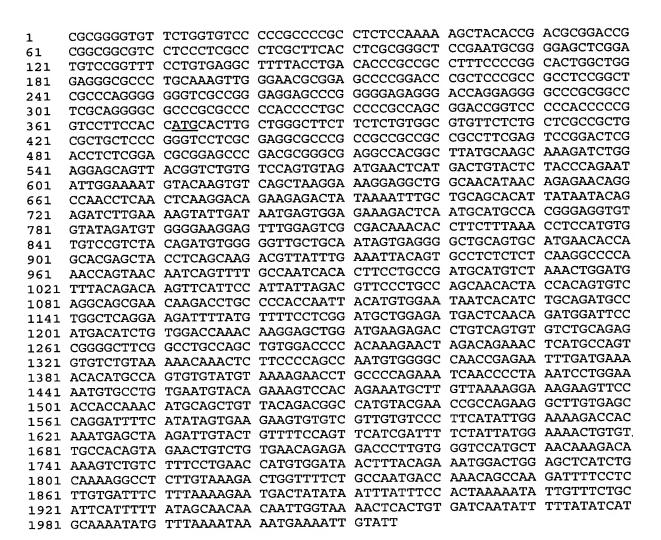


FIGURE 3A

MHLLGFFSVACSLLAAALLPGPREAPAAAAAFESGLDLSDAEPDAGEATAYASKDLEEQLRSVSSVDELM TVLYPEYWKMYKCQLRKGGWQHNREQANLNSRTEETIKFAAAHYNTEILKSIDNEWRKTQCMPREVCIDV GKEFGVATNTFFKPPCVSVYRCGGCCNSEGLQCMNTSTSYLSKTLFEITVPLSQGPKPVTISFANHTSCR CMSKLDVYRQVHSIIRRSLPATLPQCQAANKTCPTNYMWNNHICRCLAQEDFMFSSDAGDDSTDGFHDIC GPNKELDEETCQCVCRAGLRPASCGPHKELDRNSCQCVCKNKLFPSQCGANREFDENTCQCVCKRTCPRN QPLNPGKCACECTESPQKCLLKGKKFHHQTCSCYRRPCTNRQKACEPGFSYSEEVCRCVPSYWKRPQMS

FIGURE 3B